



Polymerase study: Improved detection of Salmonella and Campylobacter through the optimized use of DNA polymerases in diagnostic real-time PCR

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Aim

Introduction

Screening:

DNA Polymerase or Master mix	Performance on <i>Salmonella</i>			Performance on <i>Campylobacter</i>			Con- clusion	Price [USD/μl]	
	Rating	LoD [μg/ml]	Max dR	Rating	LoD [μg/ml]	Max dR			
Tth DNA Polymerase (Roche)	+++	1.2 × 10 ⁻⁶	42602	+++	6.6 × 10 ⁻⁷	48971	+++	1.22	
VeriQuest™ Probe qPCR Master Mix (Affymetrix)	+++	1.2 × 10 ⁻⁵	38063	+++	6.6 × 10 ⁻⁷	43067	+++	0.92	
AmpliTaQ Gold® (Applied Biosystems)	+++	1.2 × 10 ⁻⁶	34546	++	6.6 × 10 ⁻⁷	28147	+++	0.75	
HotMaster® Taq DNA Polymerase (5 Prime)	+++	1.2 × 10 ⁻⁵	39244	++	6.6 × 10 ⁻⁷	16543	+++	0.66	
TaKaRa Ex Taq® Hot Start Version (TaKaRa Bio Inc)	+++	1.2 × 10 ⁻⁵	37222	++	6.6 × 10 ⁻⁷	24549	+++	1.04	
AmpliTaQ® DNA Polymerase (Applied Biosystems)	++	1.2 × 10 ⁻⁵	40053	++	6.6 × 10 ⁻⁷	24930	++	0.80	
AmpliTaQ® 360 DNA Polymerase (Applied Biosystems)	++	1.2 × 10 ⁻⁵	39484	++	6.6 × 10 ⁻⁷	35547	++	0.56	
AmpliTaQ Gold® 360 DNA Polymerase (Applied Biosystems)	+++	1.2 × 10 ⁻⁵	41110	++	6.6 × 10 ⁻⁷	28090	++	1.29	
TaqMan® Fast Advanced Master Mix (Applied Biosystems)	++	1.2 × 10 ⁻⁶	34546	++	6.6 × 10 ⁻⁶	28331	++	1.20	
SG qPCR Master Mix (EURx)	+++	1.2 × 10 ⁻⁶	30894	+	6.6 × 10 ⁻⁶	6996	IC	0.46	
HotStarTaq® Master Mix kit (Qiagen)	+++	1.2 × 10 ⁻⁵	37751	+	6.6 × 10 ⁻⁴	14648	IC	0.99	
PicoMaxx High Fidelity PCR System (Agilent Technologies)	+	1.2 × 10 ⁻³	24797	++	6.6 × 10 ⁻⁶	19700	IC	1.00	
FastStart Taq DNA Polymerase (Roche)	+++	1.2 × 10 ⁻⁵	39644	-	6.6 × 10 ⁻²	5091	IC	1.07	
MyTaq™ HS DNA polymerase (Bioline)	+	1.2 × 10 ⁻⁵	5448	+	6.6 × 10 ⁻⁷	5338	+	0.66	
MyTaq™ DNA Polymerase (Bioline)	+	1.2 × 10 ⁻⁵	5126	+	6.6 × 10 ⁻⁶	5733	+	0.33	
Titanium™ Taq DNA Polymerase (Clontech)	-	1.2 × 10 ⁻⁵	688	+ (72°C)		6.6 × 10 ⁻⁶	8584	-	2.68
OneTaq® DNA Polymerase (New England BioLabs)	-	ND		+	6.6 × 10 ⁻⁵	12860	-	0.20	
Phusion® High-Fidelity DNA Polymerase with GC buffer (New England BioLabs)	-	1.2 × 10 ⁻⁵	2676	-	ND		-	0.84	
Pfu DNA Polymerase (Fermentas)	-	1.2	2358	-	ND		-	0.68	
Herculase II Fusion DNA Polymerase (Agilent Technologies)	-	ND		-	ND		-	0.07	

Conclusions

- ### Further evaluation of top 5:

A. Meat artificially contaminated with *Salmonella* and
B. Feces artificially contaminated with *Campylobacter*.

A.	Magnetic beads based DNA extraction			Lysis by boiling			Non-extracted		
DNA Polymerase or master mix	Rating	LoD [CFU/ml]	Max dR	Rating	LoD [CFU/ml]	Max dR	Rating	LoD [CFU/ml]	Max dR
Th	++	10 ²	59615	+	10 ³	47634	+	10 ⁵	11710
VeriQuest MM	++	10 ²	28736	++	10 ²	17216	++	10 ⁴	7430
AmpliTaq Gold	++	10 ²	64637	++	10 ²	147690	++	10 ⁴	12874
HotMaster Taq	++	10 ²	35691	+++	10 ²	97583	++	10 ⁴	23048
TaKaRa ExTaq HS	-	10 ³	27058	-	10 ⁴	47569	-	ND	257

B. DNA Polymerase or master mix	Qiagen kit extraction			Magnetic beads based DNA extraction			Lysis by boiling		
	Rating	LoD [CFU/ml]	Max dR	Rating	LoD [CFU/ml]	Max dR	Rating	LoD [CFU/ml]	Max dR
Tth	++	10 ³	22022	-	NA		-	10 ⁶	SD
VeriQuest MM	++	10 ²	12661	-	NA		+	10 ⁴	5578
AmpliTaq Gold	+	10 ³	6643	-	NA		-	NA	
HotMaster Taq	+	10 ³	9088	-	NA		-	10 ⁶	SD
TaKaRa ExTaq HS	++	10 ²	12661	+	10 ³	7861	-	10 ⁶	SD

+++ Very good, ++ Good, + Intermediate, - Poor, NA no amplification, SD single detection, *LoD could be lower, but only 10^2 to 10^6 CFU/ml was tested

Materials and Methods

16 commercially available DNA polymerases and 4 master mixes were included (see Table 1). These were evaluated on a dilution series of purified *Salmonella* ser. Typhimurium and *Campylobacter jejuni* DNA, analyzed by standardized real-time PCR assays^{2,3} using the accompanying PCR buffers for each polymerase.

The 5 best performing polymerases/kits were further evaluated using minced pork meat samples (diluted in BPW 1:10 and enriched for 18 h at 37°C followed by artificial contamination with *Salmonella* ser. Typhimurium, 10^2 - 10^6 CFU/ml) and chicken feces samples (artificially contaminated with *Campylobacter jejuni*, 10^2 - 10^6 CFU/ml). DNA extraction was performed on the samples by three different methods (Magnetic beads-based (KingFisher), lysis by boiling, and non-extracted for *Salmonella* and QIAamp® Fast DNA Stool Mini Kit (Qiagen), magnetic beads-based, and lysis by boiling for *Campylobacter*) followed by real-time PCR.

Polymerases were rated based on shape of amplification curves, amplification efficiency (AE), linear range and linearity of standard curve (R^2) and max fluorescence (Max dR)

References

1. Hedman et al (2012) Pre-PCR processing strategies. In: PCR Technologies, current innovations CRC Press, 3rd Ed.
2. Löfström et al (2009) BMC Microbiol, 9(1):85.
3. Josefsen et al (2010) AEM. 76(15):5097.

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Examples of good, intermediate, and poor performance on high to low DNA purity:

Figure 1. Performance on magnetic beads-based extracted meat samples

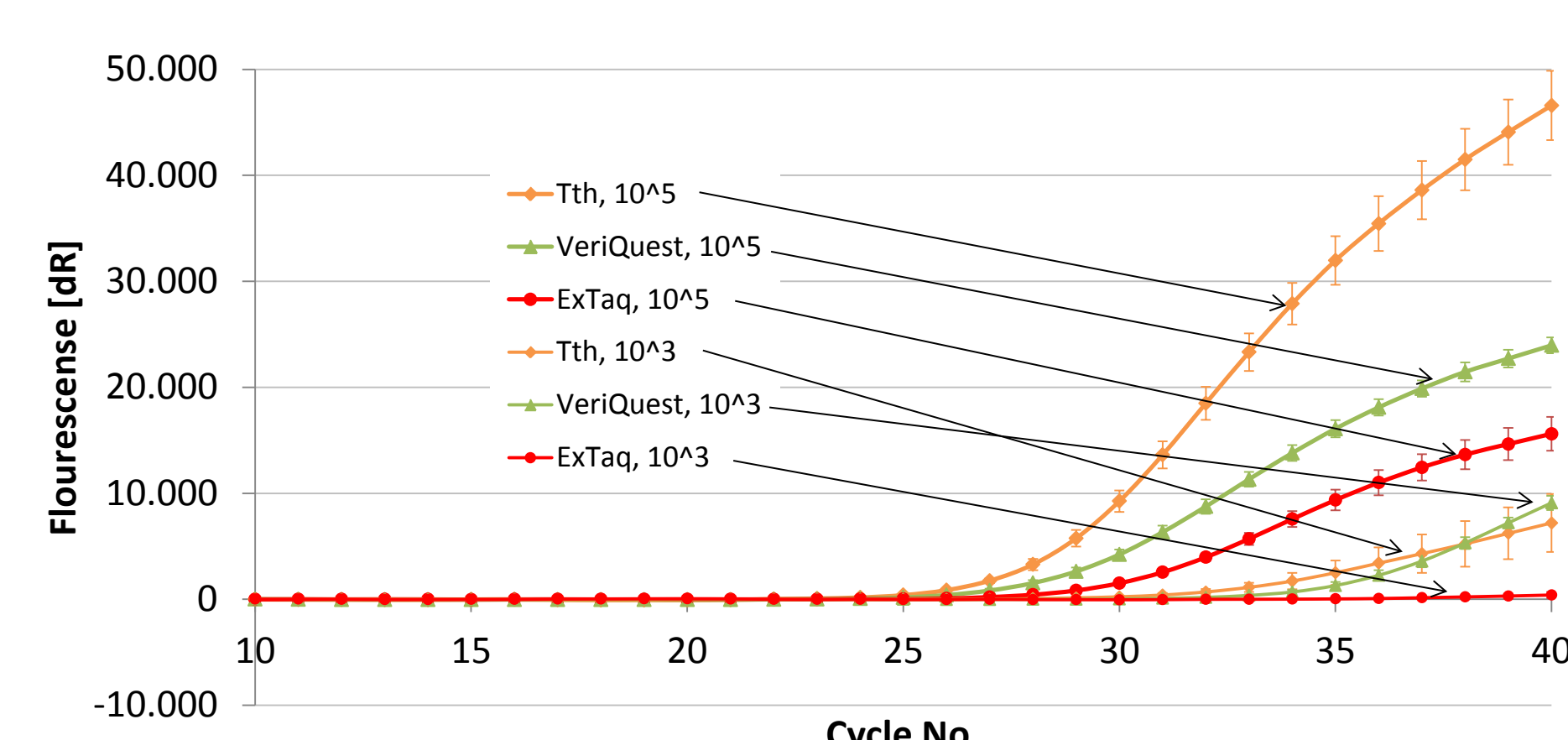


Figure 2. Performance on lysis by boiling extracted meat samples

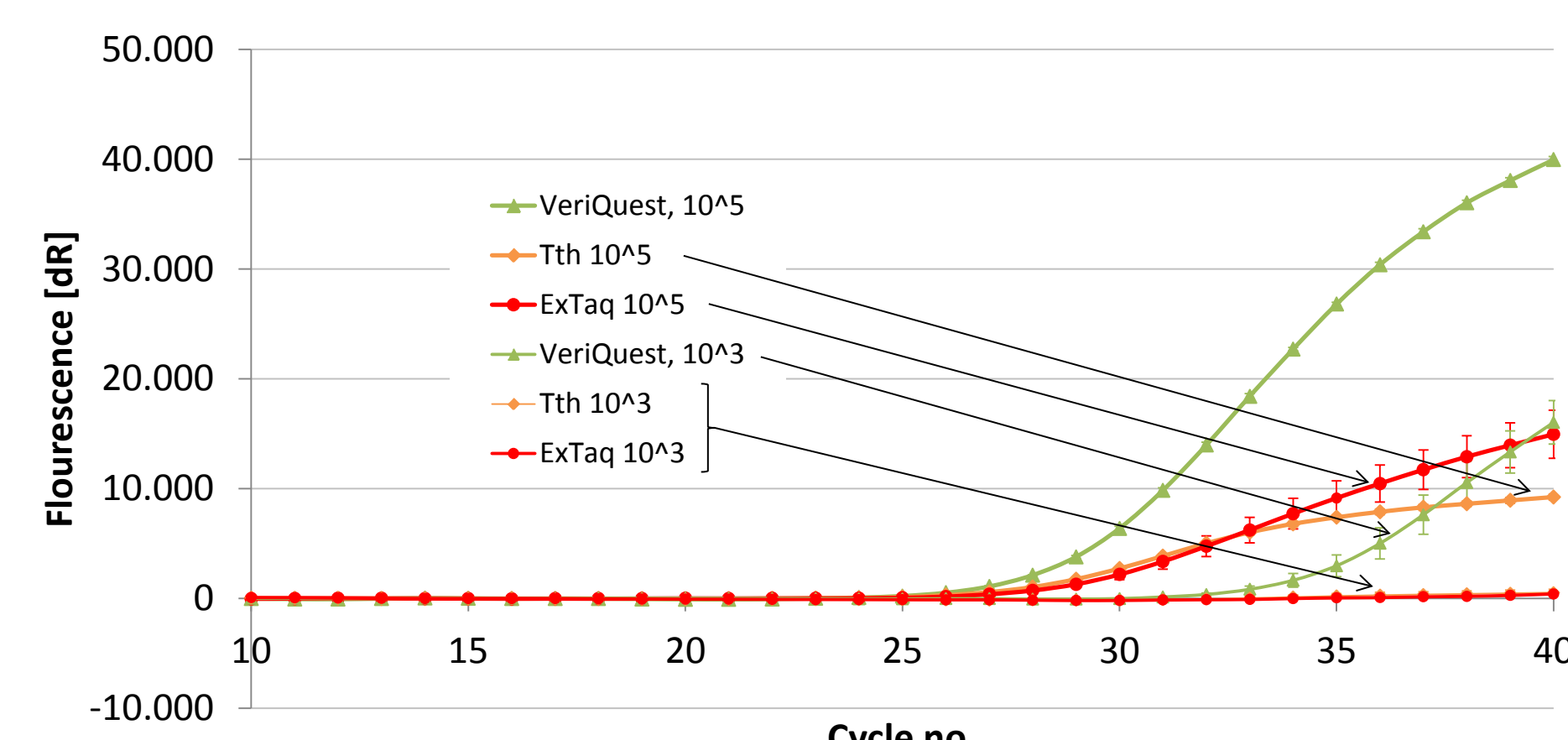
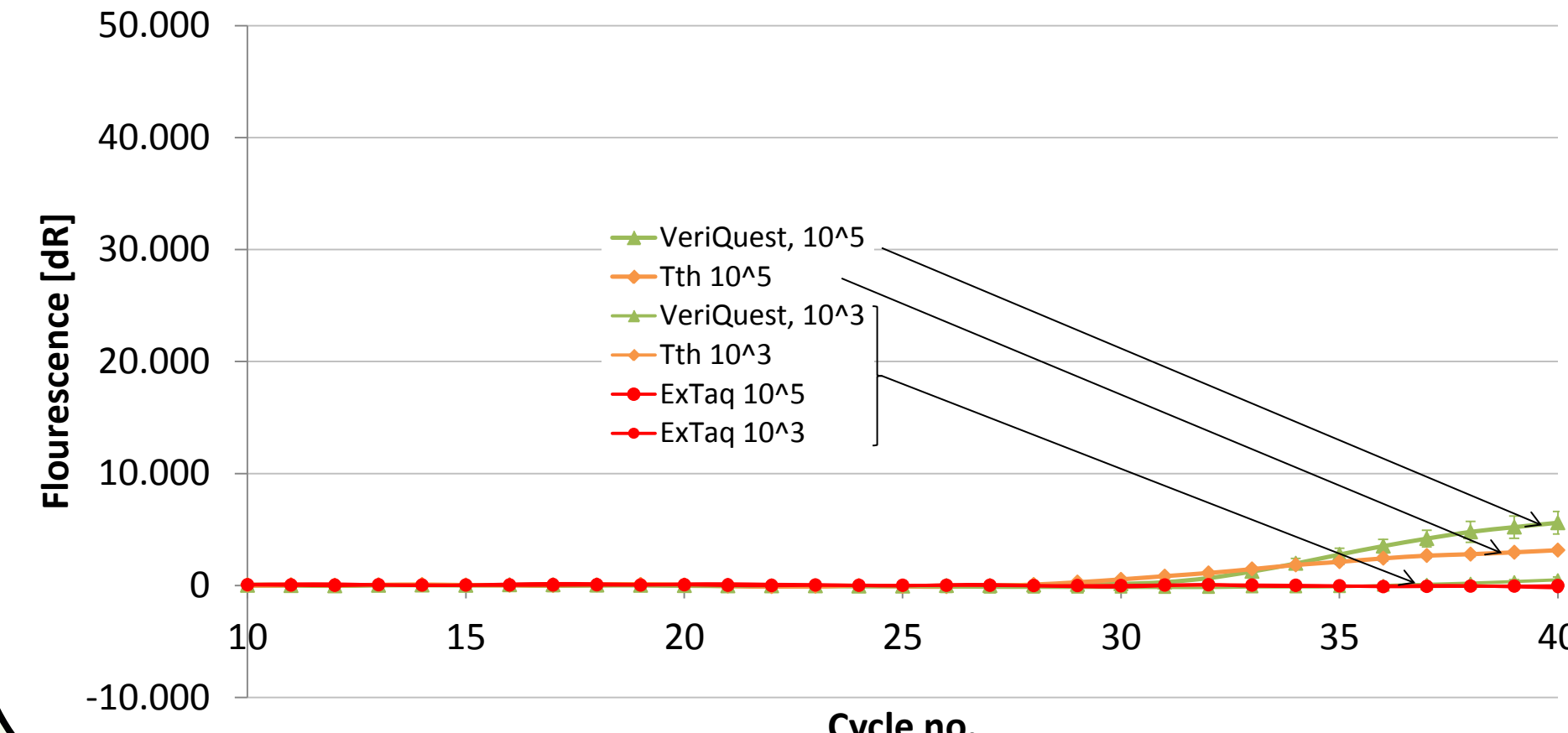


Figure 3. Performance on non-extracted meat samples



Decreasing purity of DNA extractions



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